

## Step 1:

Check that all specimens belong to the correct sample  
(if sample name is simply wrong, that will be fixed in step 2)

	er_specimen_name	er_sample_name	er_citation_names	specimen_class**	specimen_lithology**	specimen_type**
1	sr01a1	sr01a	This study			
2	sr01a2	sr01a	This study			
3	sr01c2	sr01c	This study			
4	sr01d1	sr01d	This study			
5	sr01e2	sr01e	This study			
6	sr01f2	sr01f	This study			
7	sr01g2	sr01g	This study			
8	sr01i1	sr01i	This study			